## Bioinformatics Resource Center: Interoperation

### What capability do we want?

- **■** Common repository
- **■** Common web interface
- **■** Shared datasets
  - Biodefense relevant genes
  - SNP data
  - Pathways
- **■** Tools that work on shared systems
- **■** OrthoMCL runs

#### Status: where we is.

- **■** GFF3 file format
- **≠** FTP-site



- **■** BRC-devel
- **■** Progress in some areas of data sharing GO
- **■** Progress in some areas of BRC-GO terms

#### Central Repository

- **\** Common repository for storage
- **■** Common web interface
- **■** Shared datasets
  - Biodefense relevant genes
  - SNP data
  - Pathways
- **■** Link-capability
- **■** Shared tools
- **■** Tools that work on shared systems

### Common repository, web site

- **■** Promote linkage to other centers
- **■** Promote linkage to other services
  - e.g, assembly of organism, orthoMCL, tools

#### Common web site

**♯** Promoting confusion?

# Cross BRC-linkage: URL Construction

- **■** Uniform construction for every site
  - e.g., CMR site
  - requires all sites use the same construction
- Meta site to build URL
  - Takes a datatype using key-value constructs a URL
  - Supports sites changing their web pages

#### API

- **♯** "Simple" code-base
- **■** Emphasis on reads, not writes
- **■** Emphasis on documentation
- **■** Potentially db independent
- **#** Key→value

## API: operates on what?

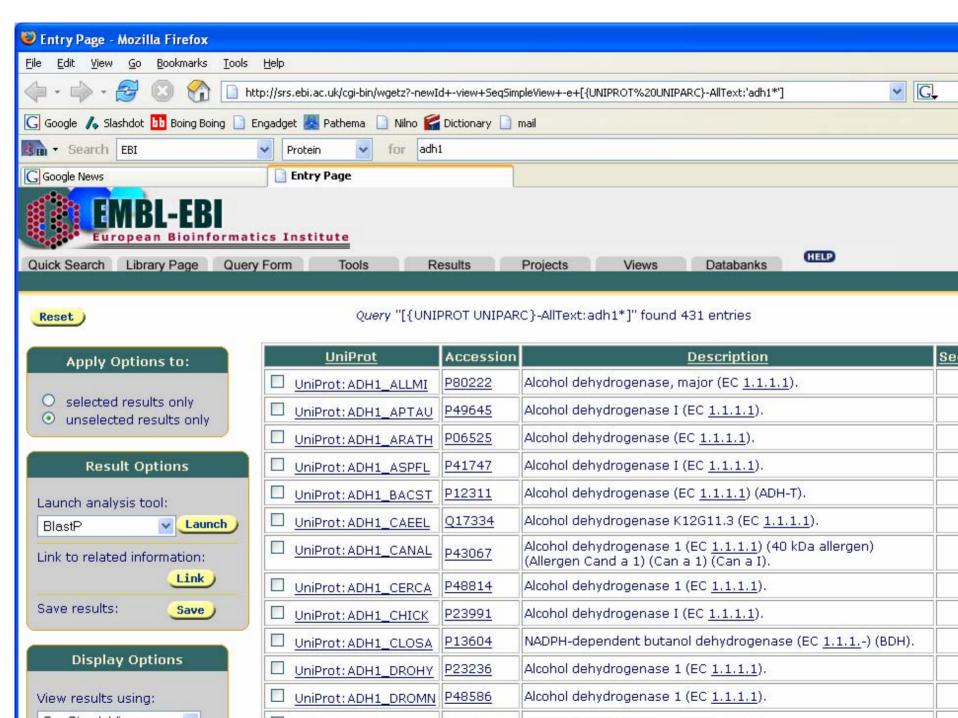
- **GFF** files:
  - Local
  - Central repository
  - Distributed
- **DB** 
  - Possible?: Gus/chado/ensembl

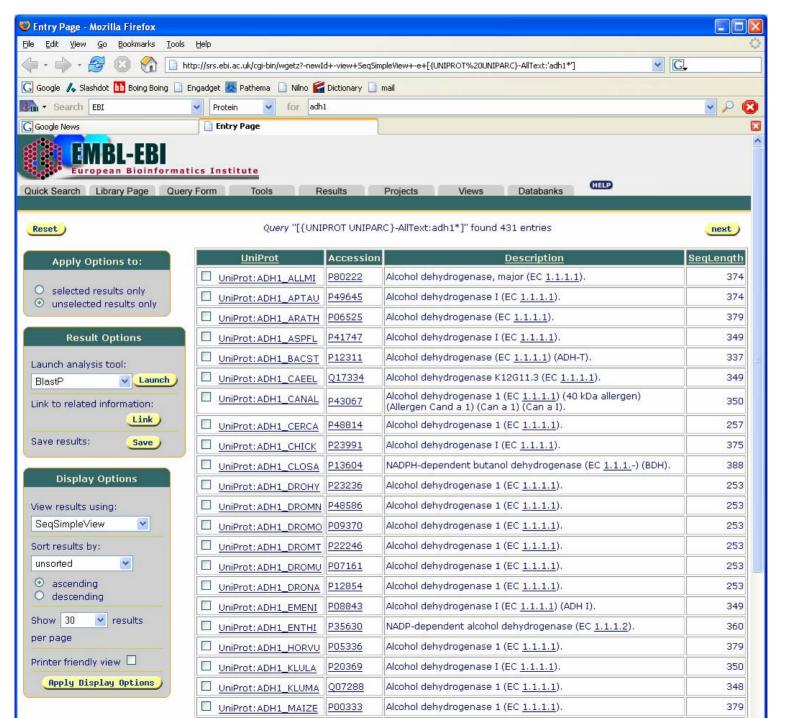
## API: coding practices

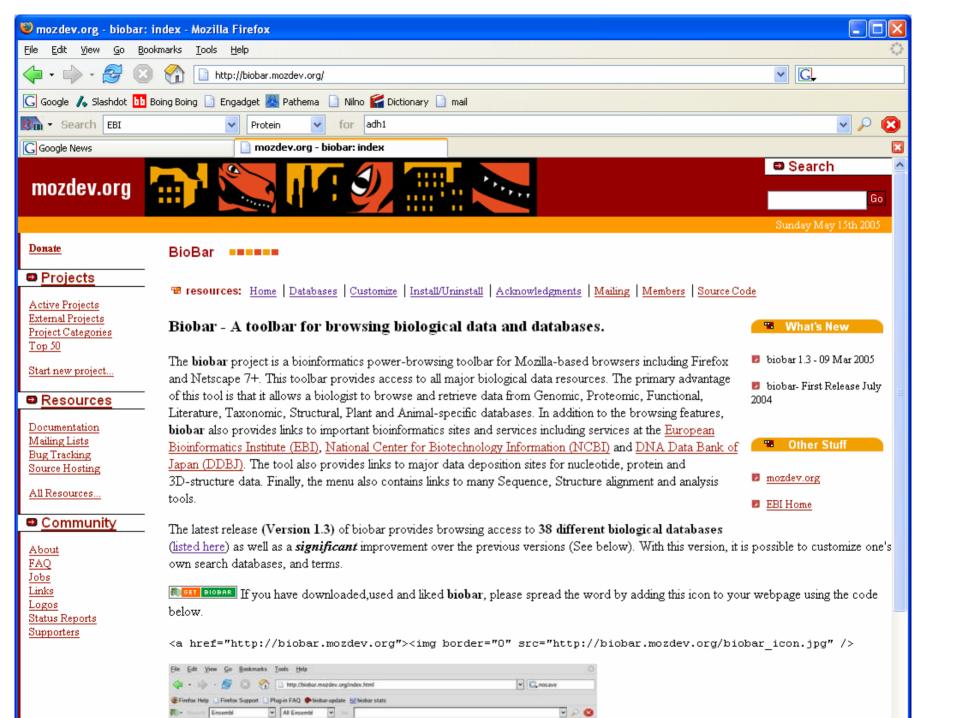
- **■** Underlying typecasting
- **■** Vigilant re-use of functions
- **♯** Rational cooperation
- **■** Version control
- **■** Documentation
- **■** Testing

#### Web Services

- **■** Note: most of GFF3 is key-value pairs
- **■** To site: key
- **■** From site: value
- **■** Potential data
  - text
  - html
  - xml
- **■** Distributed, or at central site







#### BioBar Data Sources

Data Collections	1. EBI Databases with SRS 2. NCBI Databases 3. DNA Database of Japan(DDBj)	Functional Databases	<ol> <li>aMAZE</li> <li>araCyc</li> <li>BioCyc</li> <li>KEGG</li> <li>Gene Ontology (GO)</li> <li>IntAct</li> <li>Reactome Database</li> </ol>
Sequence		Structural Databases	
Databases	1. EMBL Nucleotide Database		1. Macromolecular Structure
	2. Ensembl Genome Browser		Database (MSD)
	<ul><li>3. UniProt Sequence Database</li><li>4. InterPro</li></ul>		2. Research Collaboratory for Structural Bioinformatics
	5. Integr8		(RCSB)
	6. Expert Protein Analysis		3. Protein Data Bank Japan
	System (ExPASy)		(PDBj)
	7. Pfam		4. BMRB
	8. Alternative Splicing Database		5. OCA
	(ASD) 9. Glovar Genome Browser		6. Others
	9. Glovar Genome Browser		
<b>Animal Databases</b>	1 Coochemenson Comens	Plant Databases	1. Gramene
	1. Saccharomyces Genome Database (SGD)		2. MaizeGDB
	2. Flybase		2. MaizeGDB
	3. Wormbase		
	4. Rat Genome Database (RGD)		
	5. Mouse Genome Informatics		
	(MGI)		
	6. Zebrafish Information (ZFIN)		

# What do we want to do with our software?

- **■** Work on everyone's data?
- **■** ...with GFF3?
- **■** ... across the net?
- **■** Supply services to the BRCs?
- **■** Place on a central site?
- **■** Support as open source?

# What services do we want to supply to BRCs?

- **■** Assembly/SNP stuff
- **#** orthoMCL

#### Failure is its own reward



### What is the plan?



- **■** Central site
- **■** URL construction
- **♯** Open source software
- **API**
- **■** Web services
- **■** Biobar